

```
R version 4.2.2 (2022-10-31 ucrt) -- "Innocent and Trusting"
Copyright (C) 2022 The R Foundation for Statistical Computing
Platform: x86_64-w64-mingw32/x64 (64-bit)
```

```
R is free software and comes with ABSOLUTELY NO WARRANTY.
You are welcome to redistribute it under certain conditions.
Type 'license()' or 'licence()' for distribution details.
```

```
Natural language support but running in an English locale
```

```
R is a collaborative project with many contributors.
Type 'contributors()' for more information and
'citation()' on how to cite R or R packages in publications.
```

```
Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.
```

```
[Previously saved workspace restored]
```

```
> #Load the necessary libraries
> library(boot)
```

```
Attaching package: 'boot'
```

```
The following object is masked _by_ '.GlobalEnv':
```

```
motor
```

```
> #Load the data from CSV file
> gpa <- read.csv("C://Users//yxa210024//Desktop//Masters//spring2023//Stats for DS//mini_project
4//gpa.csv")
> #Attach the data for easy access to variables
> attach(gpa)
The following object is masked _by_ .GlobalEnv:
```

```
gpa
```

```
> #Generate scatter plots
> plot(gpa ~ act, data = gpa, pch = 20, main = "Scatterplot")
> #Calculate the Pearson correlation coefficient between gpa and act
> Cor_relation <- cor(gpa, act)
> #Define a function for bootstrap resampling
> Correlation_Resampling_function <- function(corr, indices){
+ cor(corr[indices, 1], corr[indices, 2])
+ }
> #Define a function to get the bootstrapped samples
> Corr_boots_samp <- function(corr, i = c(1:n))
+ {
+ c <- corr[i,]
+ return(c)
+ }
> #Set the number of bootstrap replicates
> Boot_replicates <- 1000
> set.seed(1000)
> #Perform bootstrap resampling for correlation between gpa and act
> Cor_boot_resample <- boot(gpa, Correlation_Resampling_function, R = Boot_replicates)
> #Display the results of bootstrap resampling
> Cor_boot_resample
```

```
ORDINARY NONPARAMETRIC BOOTSTRAP
```

```
Call:
```

```
boot(data = gpa, statistic = Correlation_Resampling_function,
      R = Boot_replicates)
```

```

Bootstrap Statistics :
      original      bias      std. error
t1* 0.2694818 0.008486976  0.1092285
> #Calculate the 95% percentile confidence interval for correlation between gpa and act
> Boot_confidence_interval(Cor_boot_resample, conf = 0.95, type = "perc")
Error in Boot_confidence_interval(Cor_boot_resample, conf = 0.95, type = "perc") :
  could not find function "Boot_confidence_interval"
> boot.ci(Cor_boot_resample, conf = 0.95, type = "perc")
BOOTSTRAP CONFIDENCE INTERVAL CALCULATIONS
Based on 1000 bootstrap replicates

```

```

CALL :
boot.ci(boot.out = Cor_boot_resample, conf = 0.95, type = "perc")

```

```

Intervals :
Level      Percentile
95%      ( 0.0575,  0.4966 )
Calculations and Intervals on Original Scale
>
>

```

```

> > Boot_replicates <- 10000
Error: unexpected '>' in ">"
> > set.seed(1000)
Error: unexpected '>' in ">"
> > #Perform bootstrap resampling for correlation between gpa and act
Error: unexpected '>' in ">"
> > Cor_boot_resample <- boot(gpa, Correlation_Resampling_function, R = Boot_replicates)
Error: unexpected '>' in ">"
> > #Display the results of bootstrap resampling
Error: unexpected '>' in ">"
> > Cor_boot_resample
Error: unexpected '>' in ">"
> Boot_replicates <- 10000
> set.seed(1000)
> Cor_boot_resample <- boot(gpa, Correlation_Resampling_function, R = Boot_replicates)
> Cor_boot_resample

```

ORDINARY NONPARAMETRIC BOOTSTRAP

```

Call:
boot(data = gpa, statistic = Correlation_Resampling_function,
      R = Boot_replicates)

```

```

Bootstrap Statistics :
      original      bias      std. error
t1* 0.2694818 0.004057848  0.1049293
> boot.ci(Cor_boot_resample, conf = 0.95, type = "perc")
BOOTSTRAP CONFIDENCE INTERVAL CALCULATIONS
Based on 10000 bootstrap replicates

```

```

CALL :
boot.ci(boot.out = Cor_boot_resample, conf = 0.95, type = "perc")

```

```

Intervals :
Level      Percentile
95%      ( 0.0701,  0.4791 )
Calculations and Intervals on Original Scale
>

```